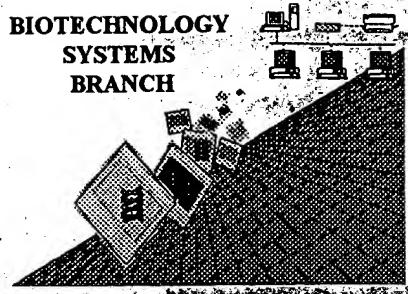


*(W. Moore)*

# RAW SEQUENCE LISTING

## ERROR REPORT

BIOTECHNOLOGY  
SYSTEMS  
BRANCH



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following CRF diskette:

Application Serial Number:

09/147,947

Art Unit / Team No. :

1652

Date Processed by STIC:

11/15/99

**THE ATTACHED PRINTOUT EXPLAINS THE ERRORS DETECTED.**

**PLEASE BE SURE TO FORWARD THIS INFORMATION TO THE APPLICANTS BY EITHER:**

**1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANTS ALONG WITH A NOTICE TO COMPLY or,**

**2) CALLING APPLICANTS AND FAXING THEM A COPY OF THE PRINTOUT WITH A NOTICE TO COMPLY**

**THIS WILL INSURE THAT THE NEXT SUBMISSION RECEIVED FROM THEM WILL BE ERROR FREE.**

**IF YOU HAVE ANY FURTHER QUESTIONS, PLEASE CALL:**

**MARK SPENCER 703-308-4212**

# Raw Sequence Listing Error Summary

## ERROR DETECTED    SUGGESTED CORRECTION

SERIAL NUMBER: 09/147,949

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1  Wrapped Nucleics The number/text at the end of each line "wrapped" down to the next line.  
This may occur if your file was retrieved in a word processor after creating it.  
Please adjust your right margin to .3, as this will prevent "wrapping".
- 2  Wrapped Aminos The amino acid number/text at the end of each line "wrapped" down to the next line.  
This may occur if your file was retrieved in a word processor after creating it.  
Please adjust your right margin to .3, as this will prevent "wrapping".
- 3  Incorrect Line Length The rules require that a line not exceed 72 characters in length. This includes spaces.
- 4  Misaligned Amino Acid Numbering The numbering under each 5th amino acid is misaligned. This may be caused by the use of tabs between the numbering. It is recommended to delete any tabs and use spacing between the numbers.
- 5  Non-ASCII This file was not saved in ASCII (DOS) text, as required by the Sequence Rules.  
Please ensure your subsequent submission is saved in ASCII text so that it can be processed.
- 6  Variable Length Sequence(s)  contain n's or Xaa's which represented more than one residue.  
As per the rules, each n or Xaa can only represent a single residue.  
Please present the maximum number of each residue having variable length and indicate in the (ix) feature section that some may be missing.
- 7  PatentIn ver. 2.0 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequence(s)  . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence.
- 8  Skipped Sequences (OLD RULES) Sequence(s)  missing. If intentional, please use the following format for each skipped sequence:  
**(2) INFORMATION FOR SEQ ID NO:X:**  
**(i) SEQUENCE CHARACTERISTICS:(Do not insert any headings under "SEQUENCE CHARACTERISTICS")**  
**(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X:**  
**This sequence is intentionally skipped**  
  
Please also adjust the "(iii) NUMBER OF SEQUENCES:" response to include the skipped sequence(s).
- 9  Skipped Sequences (NEW RULES) Sequence(s)  missing. If intentional, please use the following format for each skipped sequence.  
**<210> sequence id number**  
**<400> sequence id number**  
**000**
- 10  Use of n's or Xaa's (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.  
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.  
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 11  Use of <213>Organism (NEW RULES) Sequence(s)  are missing this mandatory field or its response.
- 12  Us of <220>Featur (NEW RULES) Sequence(s)  are missing the <220>Featur and associated headings.  
Use of <220> to <223> is MANDATORY if <213>ORGANISM is "Artificial" or "Unknown"  
Please explain source of genetic material in <220> to <223> section.  
(See "Federal Register," 6/01/98, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of new Rules)
- 13  PatentIn ver. 2.0 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other means to copy file to floppy disk.

W Moore

1652

PAGE: 1

RAW SEQUENCE LISTING  
PATENT APPLICATION US/09/147,947

DATE: 11/15/1999  
TIME: 14:07:40

Input Set: I147947.RAW

This Raw Listing contains the General Information Section and up to first 5 pages.

1 <110> APPLICANT: TSURUOKA, Nobuo  
2 YAMASHIRO, Kyoko  
3 YAMAGUCHI, Nozomi  
4 <120> TITLE OF INVENTION: Novel Serine Protease  
5 <130> FILE REFERENCE: 001560-349  
6 <140> CURRENT APPLICATION NUMBER: US/09/147,947  
7 <141> CURRENT FILING DATE: 1999-03-24  
8 <150> EARLIER APPLICATION NUMBER: PCT/JP98/03324  
9 <151> EARLIER FILING DATE: 1998-07-24  
10 <150> EARLIER APPLICATION NUMBER: JP 9/213969  
11 <151> EARLIER FILING DATE: 1997-07-24  
12 <160> NUMBER OF SEQ ID NOS: 6  
13 <170> SOFTWARE: PatentIn Ver. 2.0  
14 <210> SEQ ID NO 1  
15 <211> LENGTH: 20  
16 <212> TYPE: DNA  
17 <213> ORGANISM: Artificial Sequence  
18 <220> FEATURE:  
19 <223> OTHER INFORMATION: Synthetic DNA  
20 <400> SEQUENCE: 1 *see item 10 on Env Summary Sheet* 20  
W--> 21 gtgctca~~ng~~ chgc~~b~~caytg  
22 <210> SEQ ID NO 2  
23 <211> LENGTH: 20  
24 <212> TYPE: DNA  
25 <213> ORGANISM: Artificial Sequence  
26 <220> FEATURE:  
27 <223> OTHER INFORMATION: Synthetic DNA  
28 <400> SEQUENCE: 2 *see item 10* 20  
W--> 29 agcgg~~nn~~cc~~nc~~ cdg~~a~~rtcvcc  
30 <210> SEQ ID NO 3  
31 <211> LENGTH: 2614  
32 <212> TYPE: DNA  
33 <213> ORGANISM: Mouse  
34 <220> FEATURE:  
35 <223> OTHER INFORMATION:  
36 <400> SEQUENCE: 3  
37 cgagggtggg gttggaggtcg gactccgggc tacagagctc ctggcgctca tcgcctctgg 60  
38 ctccagcctt tgcttcgccc ggctgaccct ttgggtcccc gtgtgatcct ccagctgcc 120  
39 cgggggctgg gacagcaggg cggcggcg~~cg~~ agcgtggag ggggctctag gactctgccc 180  
40 gcccccccccc gccccc~~t~~ccg cggggacc~~cc~~ gagcc~~c~~agca tggaccacac tcggcgccgc 240  
41 agcc atg gc~~g~~ ctc gc~~c~~ cgc tgc gt~~g~~ ctg gct gt~~g~~ att tta ggg gca ctg 289  
42 Met Ala Leu Ala Arg Cys Val Leu Ala Val Ile Leu Gly Ala Leu  
43 1 5 10 15  
44 tct gta gt~~g~~ gcc cgc gct gat ccg gtc tc~~g~~ cgc tct ccc ctt cac cgc 337

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**RAW SEQUENCE LISTING**  
**PATENT APPLICATION US/09/147,947**

DATE: 11/15/1999  
 TIME: 14:07:40

Input Set: I147947.RAW

45	Ser Val Val Ala Arg Ala Asp Pro Val Ser Arg Ser Pro Leu His Arg			
46	20	25	30	
47	ccg cat ccg tcc cca ccg cgt tcc caa cac gcg cac tac ctt ccc agc	385		
48	Pro His Pro Ser Pro Pro Arg Ser Gln His Ala His Tyr Leu Pro Ser			
49	35	40	45	
50	tcg cgg cgg cca ccc agg acc ccg cgc ttc ccg ctc ccg ctg cgg atc	433		
51	Ser Arg Arg Pro Pro Arg Thr Pro Arg Phe Pro Leu Pro Leu Arg Ile			
52	50	55	60	
53	ccc gct gcc cag cgc ccg cag gtc ctc agc acc ggg cac acg ccc ccg	481		
54	Pro Ala Ala Gln Arg Pro Gln Val Leu Ser Thr Gly His Thr Pro Pro			
55	65	70	75	
56	acg att cca cgc cgc tgc ggg gca gga gag tcg tgg ggc aat gcc acc	529		
57	Thr Ile Pro Arg Arg Cys Gly Ala Gly Glu Ser Trp Gly Asn Ala Thr			
58	80	85	90	95
59	aac ctc ggc gtc ccg tgt cta cac tgg gac gag gtg ccg ccc ttc ctg	577		
60	Asn Leu Gly Val Pro Cys Leu His Trp Asp Glu Val Pro Pro Phe Leu			
61	100	105	110	
62	gag cgg tcg ccc ccg gcc agt tgg gct gag ctg cga ggg cag ccg cac	625		
63	Glu Arg Ser Pro Pro Ala Ser Trp Ala Glu Leu Arg Gly Gln Pro His			
64	115	120	125	
65	aac ttc tgc cgg agc ccg gat ggc tcg ggc aga cct tgg tgc ttc tat	673		
66	Asn Phe Cys Arg Ser Pro Asp Gly Ser Gly Arg Pro Trp Cys Phe Tyr			
67	130	135	140	
68	cgg aat gcc cag ggc aaa gta gac tgg ggc tac tgc gat tgt ggt caa	721		
69	Arg Asn Ala Gln Gly Lys Val Asp Trp Gly Tyr Cys Asp Cys Gly Gln			
70	145	150	155	
71	ggc ccg gcg ttg ccc gtc att cgc ctt gtt ggt ggg aac agt ggg cat	769		
72	Gly Pro Ala Leu Pro Val Ile Arg Leu Val Gly Gly Asn Ser Gly His			
73	160	165	170	175
74	gaa ggt cga gtg gag ctg tac cac gct ggc cag tgg ggg acc atc tgt	817		
75	Glu Gly Arg Val Glu Leu Tyr His Ala Gly Gln Trp Gly Thr Ile Cys			
76	180	185	190	
77	gac gac caa tgg gac aat gca gac gca gac gtc atc tgt agg cag ctg	865		
78	Asp Asp Gln Trp Asp Asn Ala Asp Ala Asp Val Ile Cys Arg Gln Leu			
79	195	200	205	
80	ggg ctc agt ggc att gcc aaa gca tgg cat cag gca cat ttt ggg gaa	913		
81	Gly Leu Ser Gly Ile Ala Lys Ala Trp His Gln Ala His Phe Gly Glu			
82	210	215	220	
83	gga tct ggc cca ata ttg ttg gat gaa gta cgc tgc acc gga aac gag	961		
84	Gly Ser Gly Pro Ile Leu Leu Asp Glu Val Arg Cys Thr Gly Asn Glu			
85	225	230	235	
86	ctg tca att gag caa tgt cca aag agt tcc tgg ggc gaa cat aac tgt	1009		
87	Leu Ser Ile Glu Gln Cys Pro Lys Ser Ser Trp Gly Glu His Asn Cys			
88	240	245	250	255
89	ggc cat aaa gaa gat gct gga gtg tct tgt gtt cct cta aca gat ggt	1057		
90	Gly His Lys Glu Asp Ala Gly Val Ser Cys Val Pro Leu Thr Asp Gly			
91	260	265	270	
92	gtc atc aga ctg gca gga gga aaa agt acc cat gaa ggt cgc ctg gag	1105		
93	Val Ile Arg Leu Ala Gly Gly Lys Ser Thr His Glu Gly Arg Leu Glu			
94	275	280	285	

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**RAW SEQUENCE LISTING**  
**PATENT APPLICATION US/09/147,947**

DATE: 11/15/1999  
 TIME: 14:07:40

Input Set: I147947.RAW

95	gtc tac tac aag ggg cag tgg ggg aca gtc tgt gat gat ggc tgg act	1153
96	Val Tyr Tyr Lys Gly Gln Trp Gly Thr Val Cys Asp Asp Gly Trp Thr	
97	290 295 300	
98	gag atg aac aca tac gtg gct tgt cga ctg ctg gga ttt aaa tac ggc	1201
99	Glu Met Asn Thr Tyr Val Ala Cys Arg Leu Leu Gly Phe Lys Tyr Gly	
100	305 310 315	
101	aaa cag tcc tct gtg aac cat ttt gat ggc agc aac agg ccc ata tgg	1249
102	Lys Gln Ser Ser Val Asn His Phe Asp Gly Ser Asn Arg Pro Ile Trp	
103	320 325 330 335	
104	ctg gat gac gtc agc tgc tca gga aaa gaa gtc agc ttc att cag tgt	1297
105	Leu Asp Asp Val Ser Cys Ser Gly Lys Glu Val Ser Phe Ile Gln Cys	
106	340 345 350	
107	tcc agg aga cag tgg gga agg cat gac tgc agc cat aga gaa gat gtg	1345
108	Ser Arg Arg Gln Trp Gly Arg His Asp Cys Ser His Arg Glu Asp Val	
109	355 360 365	
110	ggc ctc acc tgc tat cct gac agc gat gga cat agg ctt tct cca ggt	1393
111	Gly Leu Thr Cys Tyr Pro Asp Ser Asp Gly His Arg Leu Ser Pro Gly	
112	370 375 380	
113	ttt ccc atc aga cta gtg gat gga gag aat aag aag gaa gga cga gtg	1441
114	Phe Pro Ile Arg Leu Val Asp Gly Glu Asn Lys Lys Glu Gly Arg Val	
115	385 390 395	
116	gag gtt ttt gtc aat ggc caa tgg gga aca atc tgc gat gac gga tgg	1489
117	Glu Val Phe Val Asn Gly Gln Trp Gly Thr Ile Cys Asp Asp Gly Trp	
118	400 405 410 415	
119	acc gat aag cat gca gct gtg atc tgc cgg cag ctt ggc tat aag ggt	1537
120	Thr Asp Lys His Ala Ala Val Ile Cys Arg Gln Leu Gly Tyr Lys Gly	
121	420 425 430	
122	cct gcc aga gca agg act atg gct tat ttt ggg gaa gga aaa ggc ccc	1585
123	Pro Ala Arg Ala Arg Thr Met Ala Tyr Phe Gly Glu Gly Lys Gly Pro	
124	435 440 445	
125	atc cac atg gat aat gtg aag tgc aca gga aat gag aag gcc ctg gct	1633
126	Ile His Met Asp Asn Val Lys Cys Thr Gly Asn Glu Lys Ala Leu Ala	
127	450 455 460	
128	gac tgt gtc aaa caa gac att gga agg cac aac tgc cgc cac agt gag	1681
129	Asp Cys Val Lys Gln Asp Ile Gly Arg His Asn Cys Arg His Ser Glu	
130	465 470 475	
131	gat gca gga gtc atc tgt gac tat tta gag aag aaa gca tca agt agt	1729
132	Asp Ala Gly Val Ile Cys Asp Tyr Leu Glu Lys Lys Ala Ser Ser Ser	
133	480 485 490 495	
134	ggt aat aaa gag atg ctc tca tct gga tgt gga ctg agg tta ctg cac	1777
135	Gly Asn Lys Glu Met Leu Ser Ser Gly Cys Gly Leu Arg Leu Leu His	
136	500 505 510	
137	cgt cgg cag aaa cgg atc att ggt ggg aac aat tct tta agg ggt gcc	1825
138	Arg Arg Gln Lys Arg Ile Ile Gly Gly Asn Asn Ser Leu Arg Gly Ala	
139	515 520 525	
140	tgg cct tgg cag gct tcc ctc agg ctg agg tcg gcc cat gga gac ggc	1873
141	Trp Pro Trp Gln Ala Ser Leu Arg Leu Arg Ser Ala His Gly Asp Gly	
142	530 535 540	
143	agg ctg ctt tgt gga gct acc ctt ctg agt agc tgc tgg gtc ctg aca	1921
144	Arg Leu Leu Cys Gly Ala Thr Leu Leu Ser Ser Cys Trp Val Leu Thr	

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**RAW SEQUENCE LISTING**  
**PATENT APPLICATION US/09/147,947**

DATE: 11/15/1999  
 TIME: 14:07:40

Input Set: I147947.RAW

145	545	550	555	
146	gct gca cac tgc ttc aaa agg tac gga aac aac tcg agg agc tat gca			1969
147	Ala Ala His Cys Phe Lys Arg Tyr Gly Asn Asn Ser Arg Ser Tyr Ala			
148	560	565	570	575
149	gtt cga gtt ggg gat tat cat act ctg gta cca gag gag ttt gaa caa			2017
150	Val Arg Val Gly Asp Tyr His Thr Leu Val Pro Glu Glu Phe Glu Gln			
151	580	585	590	
152	gaa ata ggg gtt caa cag att gtg att cac agg aac tac agg cca gac			2065
153	Glu Ile Gly Val Gln Gln Ile Val Ile His Arg Asn Tyr Arg Pro Asp			
154	595	600	605	
155	aga agc gac tat gac att gcc ctg gtt aga ttg caa gga cca ggg gag			2113
156	Arg Ser Asp Tyr Asp Ile Ala Leu Val Arg Leu Gln Gly Pro Gly Glu			
157	610	615	620	
158	caa tgt gcc aga cta agc acc cac gtt ttg cca gcc tgt tta cct cta			2161
159	Gln Cys Ala Arg Leu Ser Thr His Val Leu Pro Ala Cys Leu Pro Leu			
160	625	630	635	
161	tgg aga gag agg cca cag aaa aca gcc tcc aac tgt cac ata aca gga			2209
162	Trp Arg Glu Arg Pro Gln Lys Thr Ala Ser Asn Cys His Ile Thr Gly			
163	640	645	650	655
164	tgg gga gac aca ggt cgt gcc tac tca aga act cta caa caa gct gct			2257
165	Trp Gly Asp Thr Gly Arg Ala Tyr Ser Arg Thr Leu Gln Gln Ala Ala			
166	660	665	670	
167	gtg cct ctg tta ccc aag agg ttt tgt aaa gag agg tac aag gga cta			2305
168	Val Pro Leu Leu Pro Lys Arg Phe Cys Lys Glu Arg Tyr Lys Gly Leu			
169	675	680	685	
170	ttt act ggg aga atg ctc tgt gct ggg aac ctc caa gaa gac aac cgt			2353
171	Phe Thr Gly Arg Met Leu Cys Ala Gly Asn Leu Gln Glu Asp Asn Arg			
172	690	695	700	
173	gtg gac agc tgc cag gga gac agt gga gga cca ctc atg tgt gaa aag			2401
174	Val Asp Ser Cys Gln Gly Asp Ser Gly Gly Pro Leu Met Cys Glu Lys			
175	705	710	715	
176	cct gat gag tcc tgg gtt gtg tat ggg gtg act tcc tgg ggg tat gga			2449
177	Pro Asp Glu Ser Trp Val Val Tyr Gly Val Thr Ser Trp Gly Tyr Gly			
178	720	725	730	735
179	tgt gga gtc aaa gac act cct gga gtt tat acc aga gtc ccc gcc ttt			2497
180	Cys Gly Val Lys Asp Thr Pro Gly Val Tyr Thr Arg Val Pro Ala Phe			
181	740	745	750	
182	gta cct tgg ata aaa agt gtc acc agt ctg taacttatgg aaagctcaag			2547
183	Val Pro Trp Ile Lys Ser Val Thr Ser Leu			
184	755	760		
185	aaaaatagtaa aacagtaacc attcagtctt catacttggc accatgccag aaaaaaaaaa			2607
186	aaaaaaaa			2614
187	<210> SEQ ID NO 4			
188	<211> LENGTH: 761			
189	<212> TYPE: PRT			
190	<213> ORGANISM: Mouse			
191	<220> FEATURE:			
192	<223> OTHER INFORMATION:			
193	<400> SEQUENCE: 4			
194	Met Ala Leu Ala Arg Cys Val Leu Ala Val Ile Leu Gly Ala Leu Ser			

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**RAW SEQUENCE LISTING**  
**PATENT APPLICATION US/09/147,947**

 DATE: 11/15/1999  
 TIME: 14:07:40

Input Set: I147947.RAW

195	1	5	10	15
196	Val Val Ala Arg Ala Asp Pro Val Ser Arg Ser Pro Leu His Arg Pro			
197	20	25	30	
198	His Pro Ser Pro Pro Arg Ser Gln His Ala His Tyr Leu Pro Ser Ser			
199	35	40	45	
200	Arg Arg Pro Pro Arg Thr Pro Arg Phe Pro Leu Pro Leu Arg Ile Pro			
201	50	55	60	
202	Ala Ala Gln Arg Pro Gln Val Leu Ser Thr Gly His Thr Pro Pro Thr			
203	65	70	75	80
204	Ile Pro Arg Arg Cys Gly Ala Gly Glu Ser Trp Gly Asn Ala Thr Asn			
205	85	90	95	
206	Leu Gly Val Pro Cys Leu His Trp Asp Glu Val Pro Pro Phe Leu Glu			
207	100	105	110	
208	Arg Ser Pro Pro Ala Ser Trp Ala Glu Leu Arg Gly Gln Pro His Asn			
209	115	120	125	
210	Phe Cys Arg Ser Pro Asp Gly Ser Gly Arg Pro Trp Cys Phe Tyr Arg			
211	130	135	140	
212	Asn Ala Gln Gly Lys Val Asp Trp Gly Tyr Cys Asp Cys Gly Gln Gly			
213	145	150	155	160
214	Pro Ala Leu Pro Val Ile Arg Leu Val Gly Gly Asn Ser Gly His Glu			
215	165	170	175	
216	Gly Arg Val Glu Leu Tyr His Ala Gly Gln Trp Gly Thr Ile Cys Asp			
217	180	185	190	
218	Asp Gln Trp Asp Asn Ala Asp Ala Asp Val Ile Cys Arg Gln Leu Gly			
219	195	200	205	
220	Leu Ser Gly Ile Ala Lys Ala Trp His Gln Ala His Phe Gly Glu Gly			
221	210	215	220	
222	Ser Gly Pro Ile Leu Leu Asp Glu Val Arg Cys Thr Gly Asn Glu Leu			
223	225	230	235	240
224	Ser Ile Glu Gln Cys Pro Lys Ser Ser Trp Gly Glu His Asn Cys Gly			
225	245	250	255	
226	His Lys Glu Asp Ala Gly Val Ser Cys Val Pro Leu Thr Asp Gly Val			
227	260	265	270	
228	Ile Arg Leu Ala Gly Gly Lys Ser Thr His Glu Gly Arg Leu Glu Val			
229	275	280	285	
230	Tyr Tyr Lys Gly Gln Trp Gly Thr Val Cys Asp Asp Gly Trp Thr Glu			
231	290	295	300	
232	Met Asn Thr Tyr Val Ala Cys Arg Leu Leu Gly Phe Lys Tyr Gly Lys			
233	305	310	315	320
234	Gln Ser Ser Val Asn His Phe Asp Gly Ser Asn Arg Pro Ile Trp Leu			
235	325	330	335	
236	Asp Asp Val Ser Cys Ser Gly Lys Glu Val Ser Phe Ile Gln Cys Ser			
237	340	345	350	
238	Arg Arg Gln Trp Gly Arg His Asp Cys Ser His Arg Glu Asp Val Gly			
239	355	360	365	
240	Leu Thr Cys Tyr Pro Asp Ser Asp Gly His Arg Leu Ser Pro Gly Phe			
241	370	375	380	
242	Pro Ile Arg Leu Val Asp Gly Glu Asn Lys Lys Glu Gly Arg Val Glu			
243	385	390	395	400
244	Val Phe Val Asn Gly Gln Trp Gly Thr Ile Cys Asp Asp Gly Trp Thr			

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**VERIFICATION SUMMARY  
PATENT APPLICATION US/09/147,947**

DATE: 11/15/1999

TIME: 14:07:40

Input Set: I147947.RAW

Line ? Error/Warning	Original Text
21 W "N" or "Xaa" used: Feature required	gtgctcacng cngcbcaytg
29 W "N" or "Xaa" used: Feature required	agcggnccnc cdgartcvcc